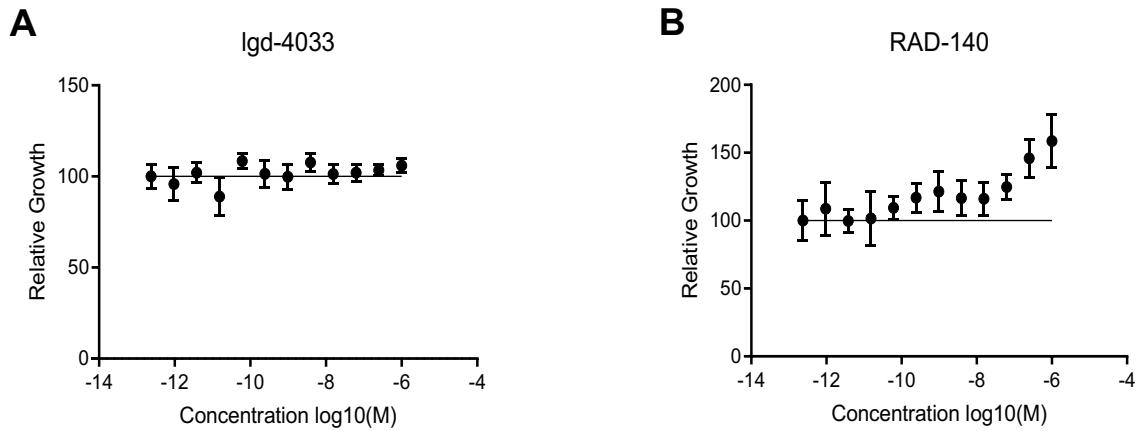


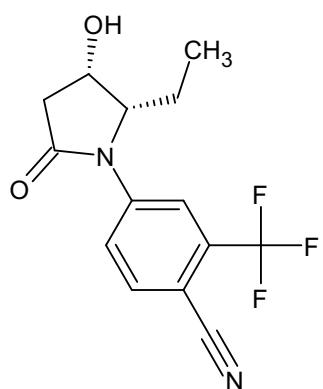
Supplemental Figure 1



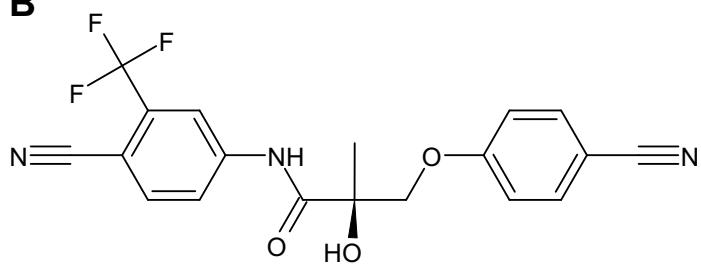
Supplemental Figure 1: SARM effects on cell growth. Dose response curves were performed on LNCaP cells of SARMs **A)** lgd-4033 and **B)** RAD-140. (n=4) (Data represent mean \pm SD).

Supplemental Figure 2

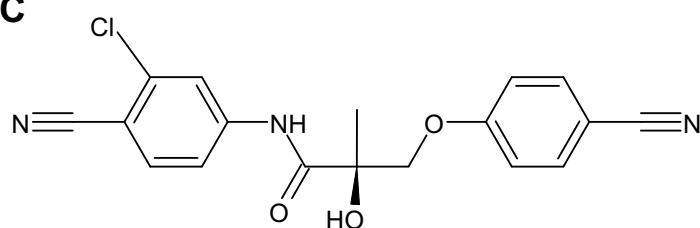
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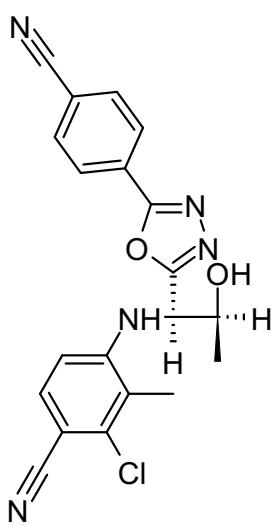
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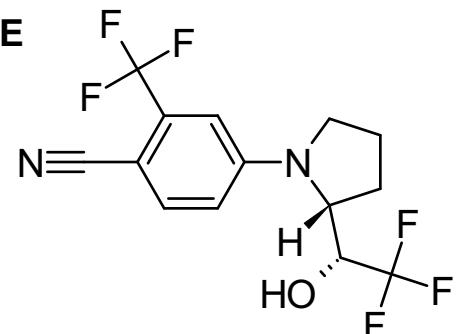
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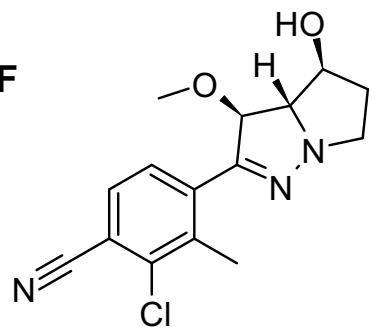
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E



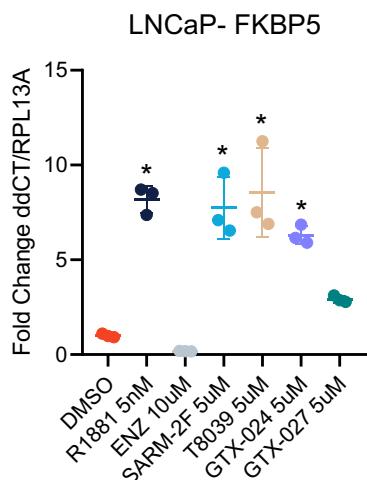
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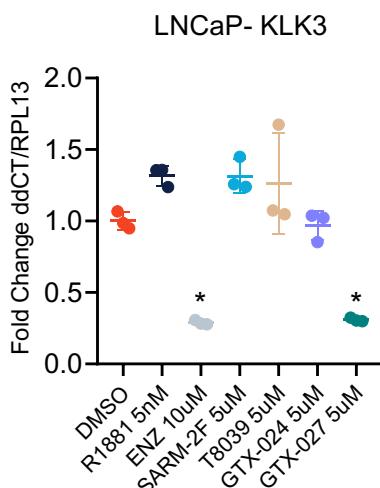
Supplemental Figure 2: Non-steroidal Selective Androgen Receptor Modulators.
Structures of **A**) SARM-2F, **B**) GTx-024, **C**) GTx-027, **D**) RAD-140, **E**) LGD-4033, and **F**) Compound-26.

Supplemental Figure 3

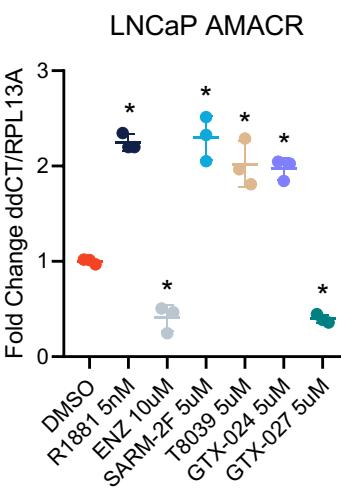
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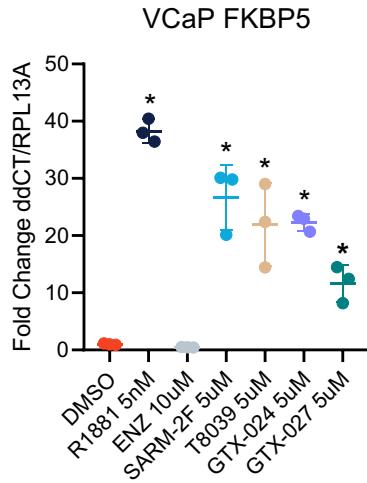
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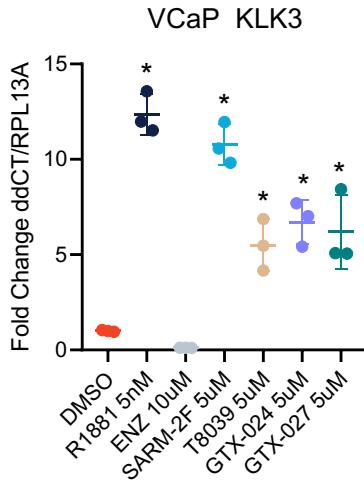
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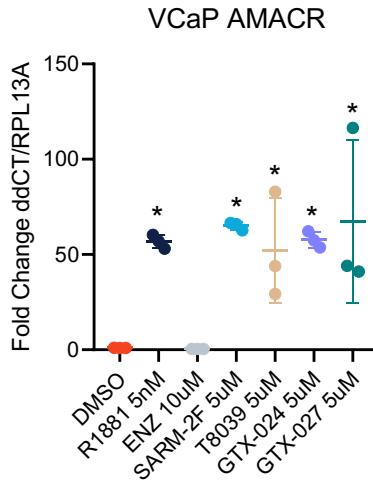
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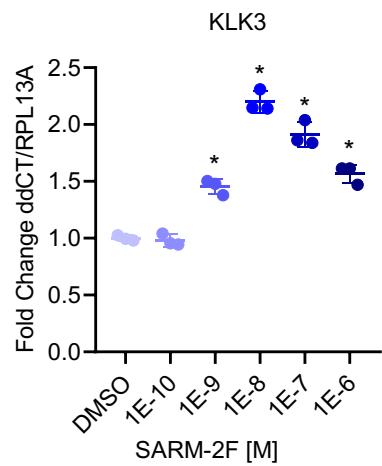
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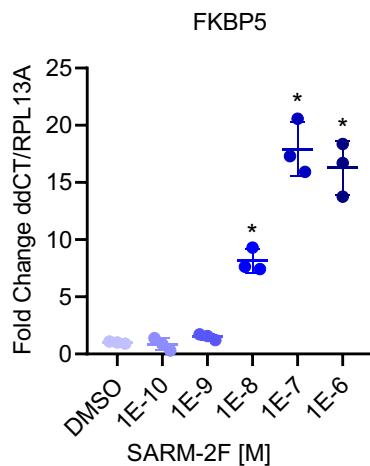
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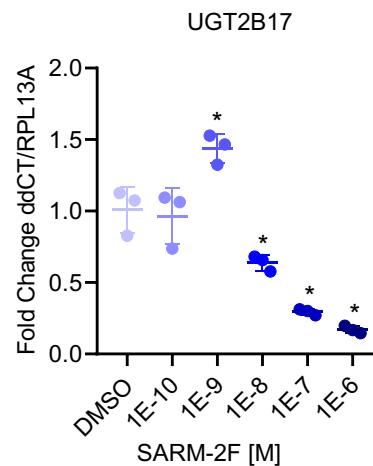
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H



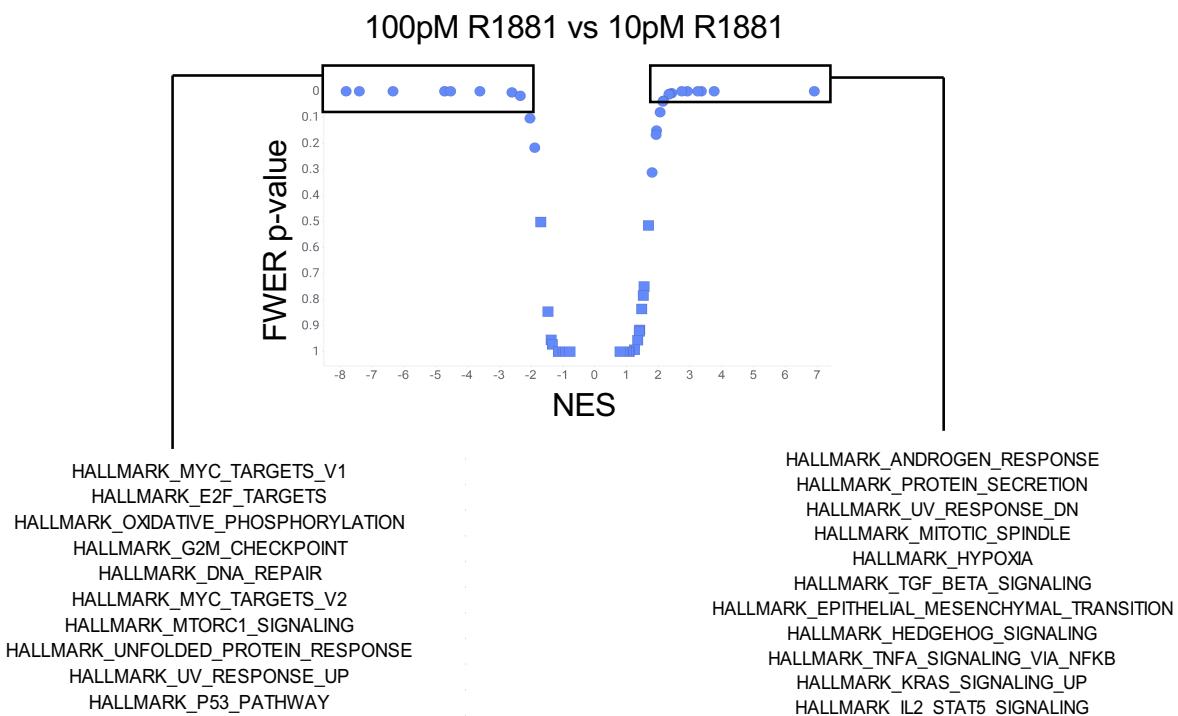
I



Supplemental Figure 3 SARM effects on gene expression. qRT-PCRs of FKBP5, KLK3 and AMACR in **A-C)** LNCaP and **D-F)** VCaP cells. (n=3) (* = p<0.05) (Data represent mean ± SD). qRT-PCRs of AR-induced genes **G) KLK**, **H) FKBP5**, and AR-repressed gene **I) UGT2B17** in LNCaP cells cultured 48hrs with a dose-range of SARM-2F. (n=3) (* = p<0.05 one-way ANOVA Dunnett's multiple comparisons test) (Data represent mean ± SD).

Supplemental Figure 4

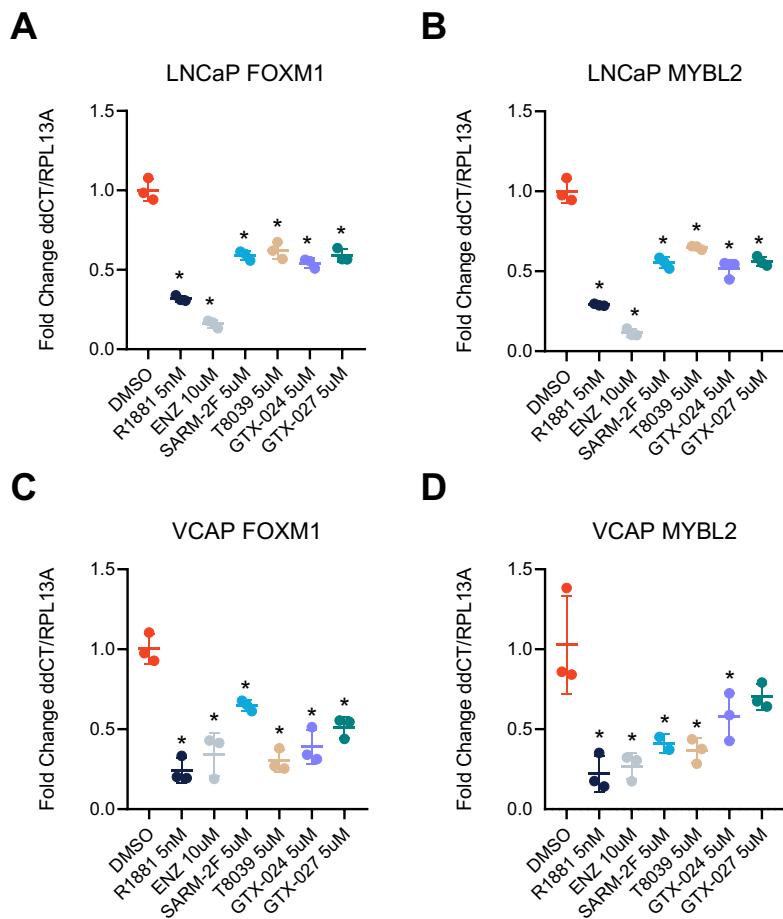
A



Supplemental Figure 4: GSEA comparison of SARMs and steroidal ligand concentrations.

A) Plot of FWER p-values vs NES scores of HALLMARK genesets for 100pM R1881 vs 10pM R1881 GSEA analyses.

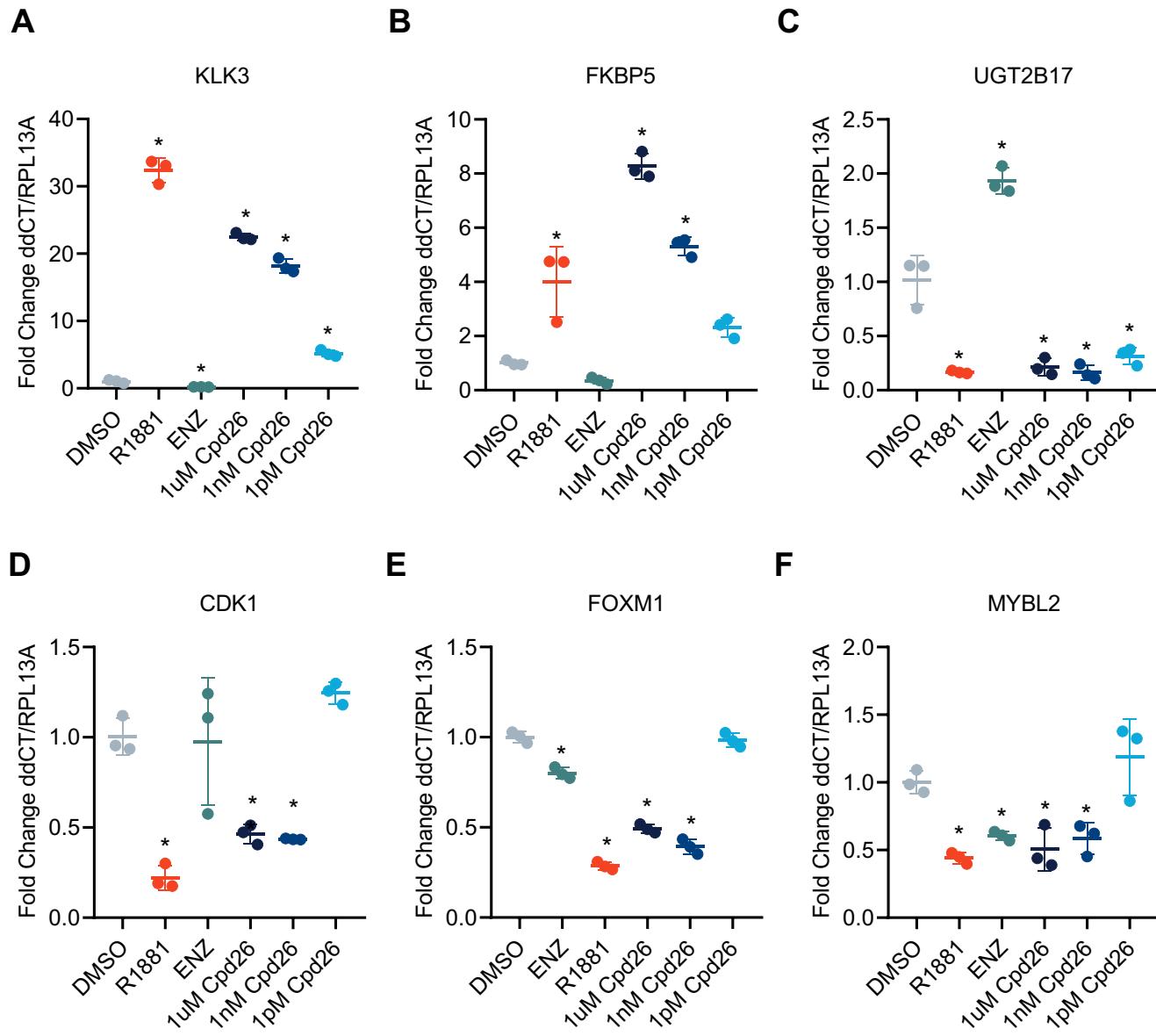
Supplemental Figure 5



Supplemental Figure 5: SARMs suppress cell-cycle gene expression.

qRT-PCR quantitation of FOXM1 and MYBL2 in **(A,B)** LNCaP and **(C,D)** VCaP cells exposed to the synthetic androgen R1881, the AR antagonist ENZ and various SARMs. (n=3) (*) = p<0.05 one-way ANOVA Dunnett's multiple comparisons test) (Data represent mean ± SD).

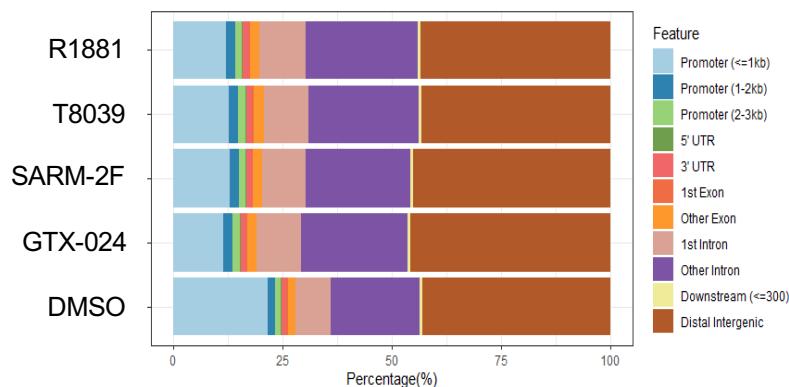
Supplemental Figure 6



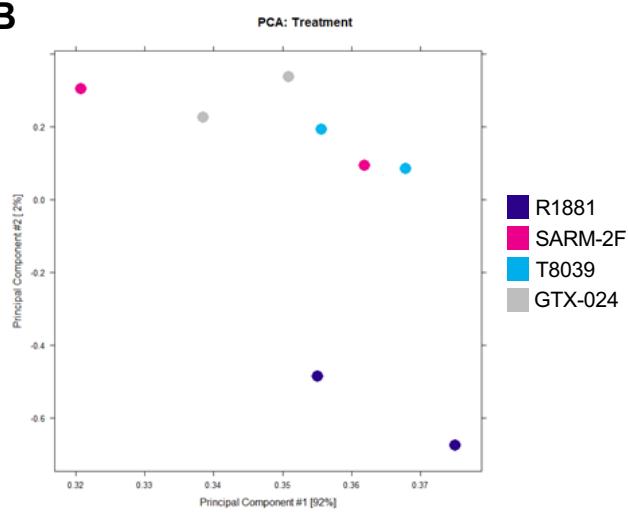
Supplemental Figure 6: SARM cpd26 induces AR-regulated gene expression changes in VCaP cells. qRT-PCR quantitation of AR-induced genes **A)** KLK3, and **B)** FKBP5, AR-repressed gene **C)** UGT2B17, and cell-cycle related genes **D)** CDK1, **E)** FOXM1 and **F)** MYBL2 in VCaP cells 48 hours after treatment. (n=3) (* = p<0.05 one-way ANOVA Dunnett's multiple comparisons test). (Data represent mean ± SD).

Supplemental Figure 7

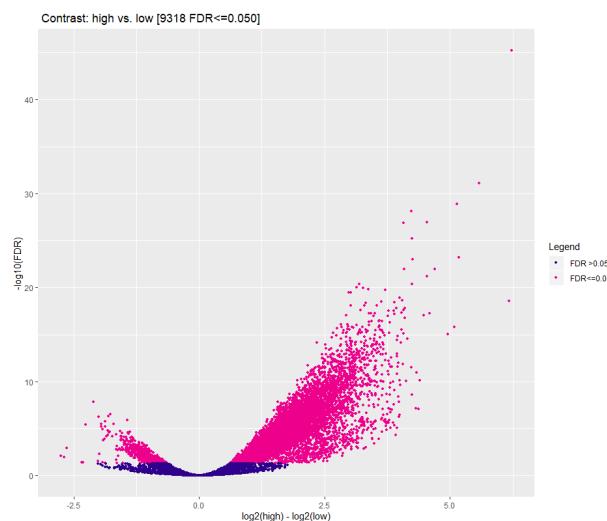
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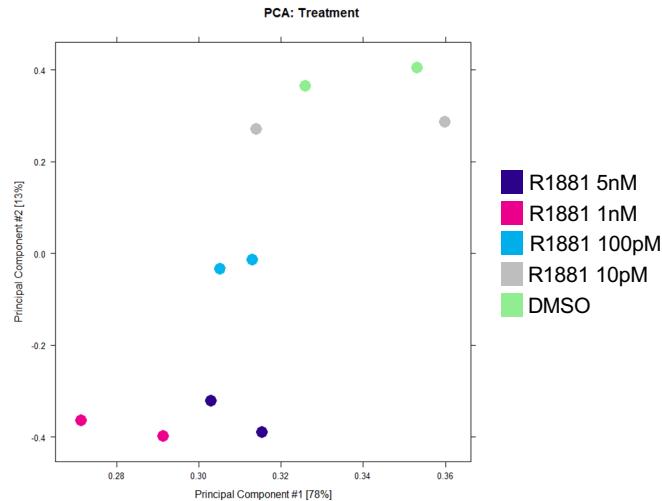
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C

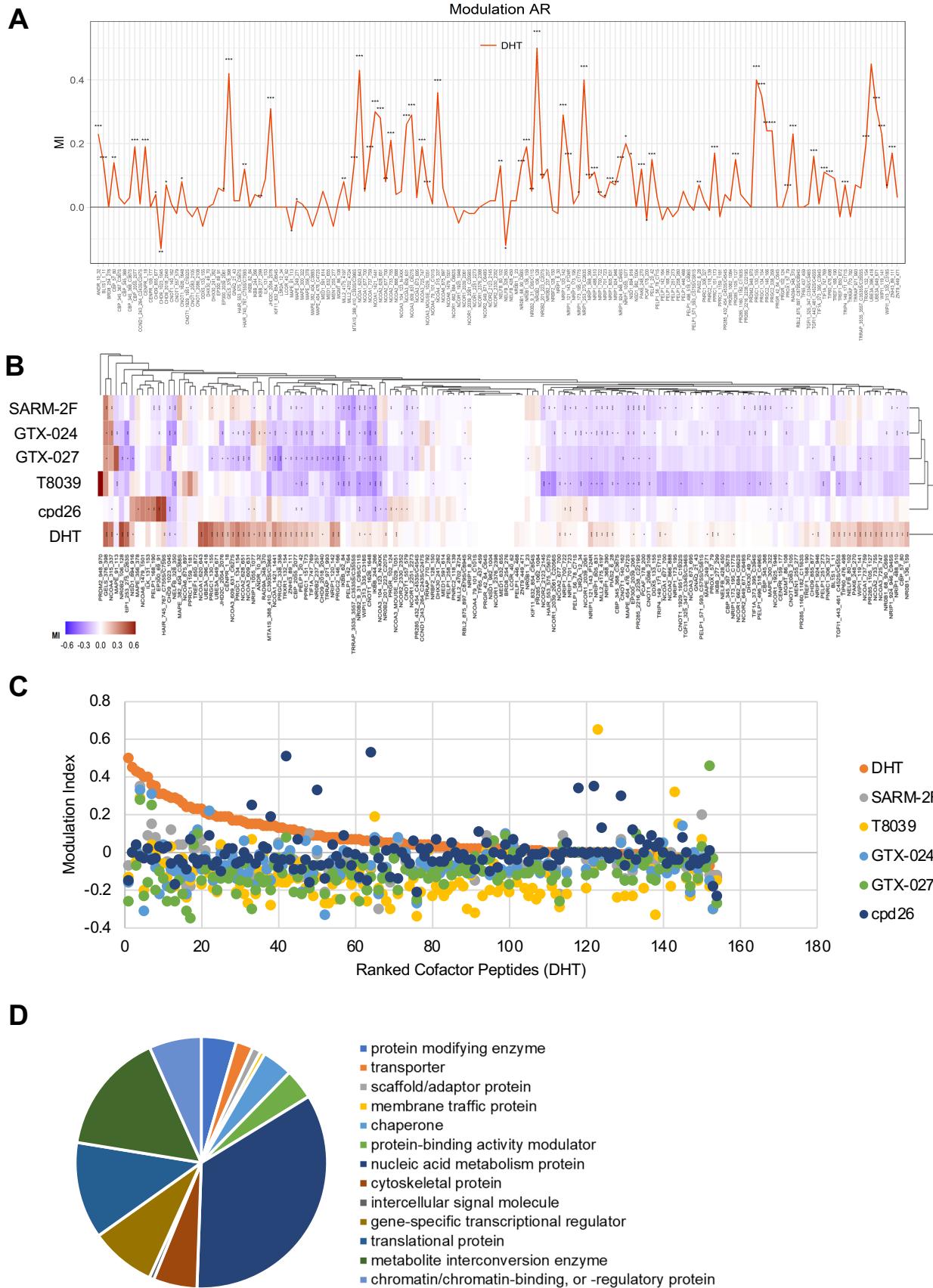


D



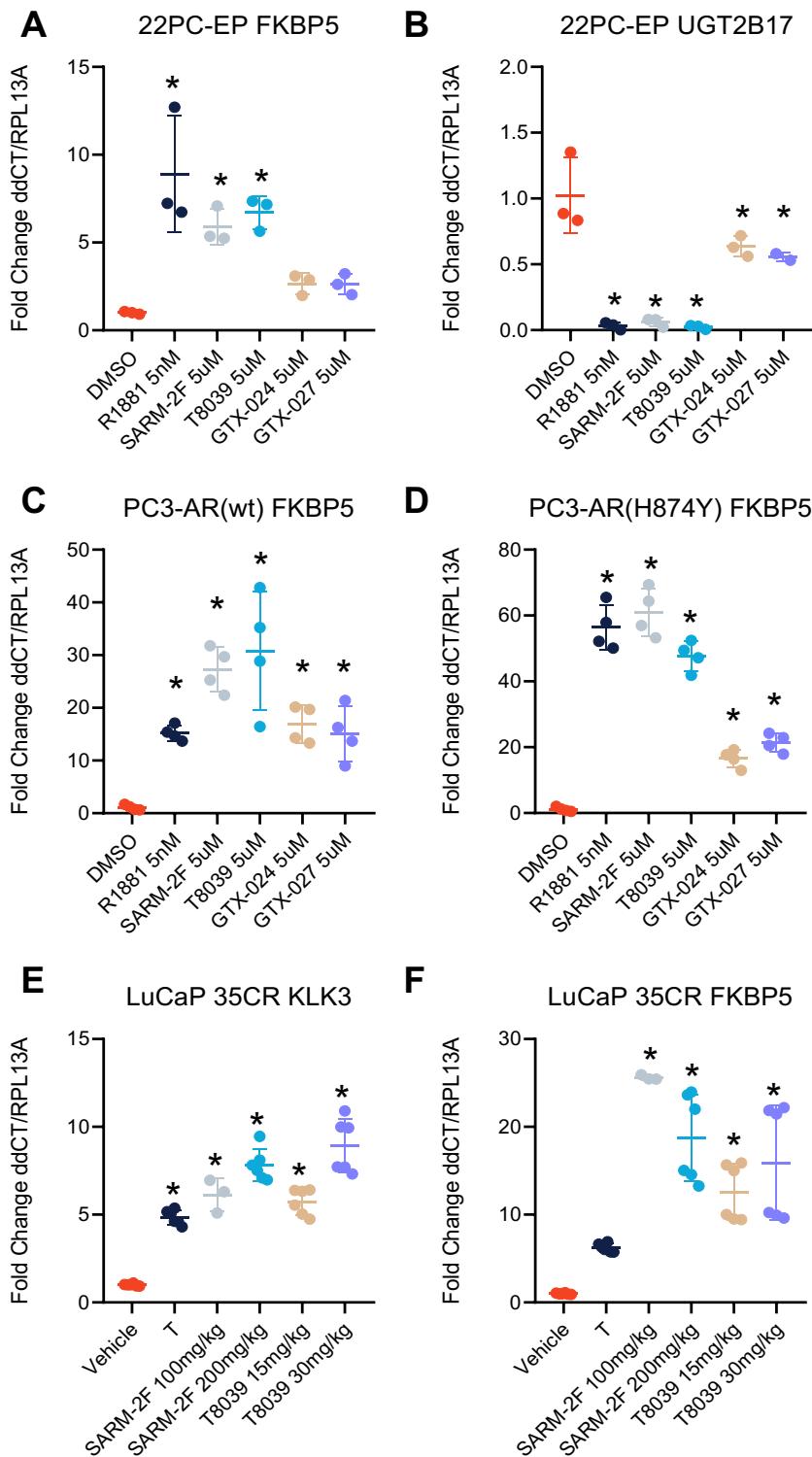
Supplemental Figure 7. Comparison of steroid and SARM induced AR cistromes using CUT&RUN. **A)** Annotation of genomic features bound by R1881 and SARM liganded AR. **B)** Principal component plot comparing the AR cistromes of SARM and high-dose R1881 treated cells. **C)** Volcano plot of differentially bound sites between low and high-dose AR sites. **D)** Principal component plot comparing AR cistromes of a dose-range of R1881 treated cells.

Supplemental Figure 8



Supplemental Figure 8. Analyses of SARM induced AR cofactor binding. **A)** MARCONI modulation Index (MI = $\log_{10}(\text{DHT signal} / \text{DMSO signal})$) values of nuclear receptor cofactors for DHT agonized AR-LBD (n=3). **B)** Heat map comparing SARM agonized and DHT agonized cofactor binding MI values. **C)** DHT-ranked plot of MARCONI MI values. **D)** Gene Ontology annotation of proteins bound to AR in both SARM-2F and R1881 samples as detected by RIME.

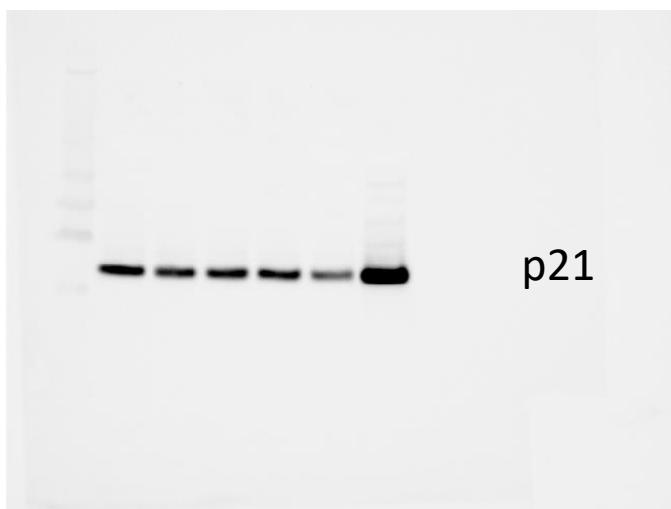
Supplemental Figure 9



Supplemental Figure 9. AR target gene expression in SARM-treated 22PC-EP, PC3-AR, and LuCAP35CR PDX. qRT-PCR analyses were performed on 22PC-EP treated for 48 hours at the indicated doses of SARMs under normal growth conditions for: **A**) AR activated gene FKBP5, and **B**) AR repressed gene UGT2B17. (n=3)(* = p < 0.05) (Data represent mean ± SD). qRT-PCR analyses were performed for FKBP5 on PC3 cells engineered to express **C**) wild-type AR or **D**) AR-H874Y and treated for 48 hours at the indicated doses of SARMs under normal growth conditions (n=4)(* = p < 0.05). qRT-PCR on RNA extracted 48 hours after PDX model LuCaP 35CR was treated with the indicated doses of SARMs, orally, or an IM injection of 40mg/kg testosterone for: **E**) KLK3 and **F**) FKBP5. Each dose is represented by two tumors analyzed separately by qRT-PCR then combined to graph (n=3+3) (* = p < 0.05 one-way ANOVA Dunnett's multiple comparisons test) (Data represent mean ± SD).

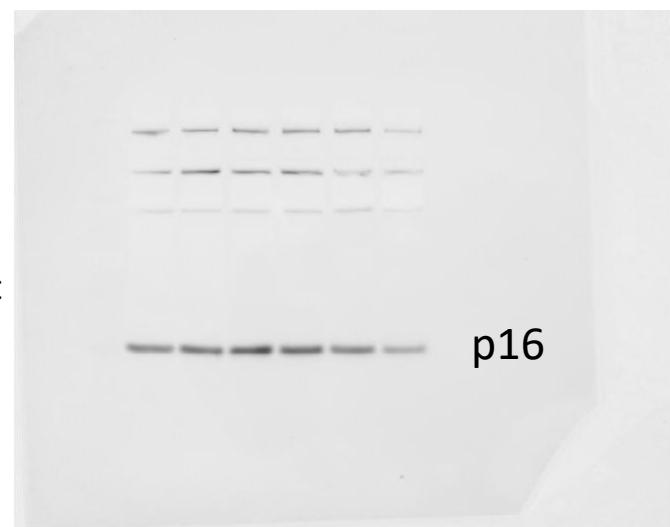
Full unedited gels for Figure 2E

1 2 3 4 5 6

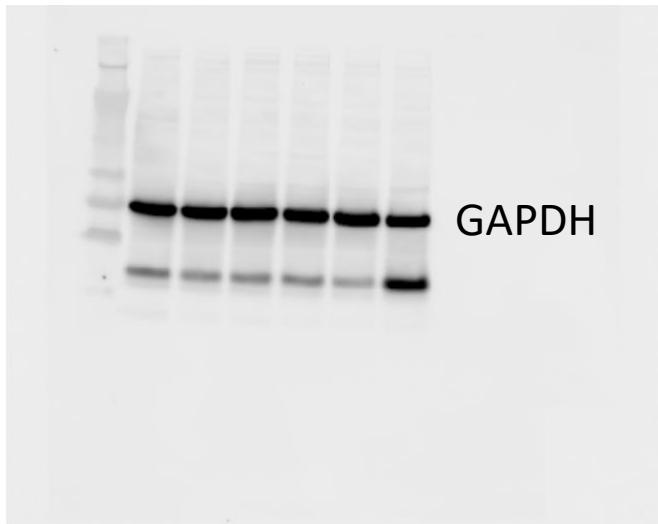


Lanes:
1) LNCaP
2) R1881
3) SARM-2F
4) T8039
5) GTx-024
6) Mitomycin-C

1 2 3 4 5 6

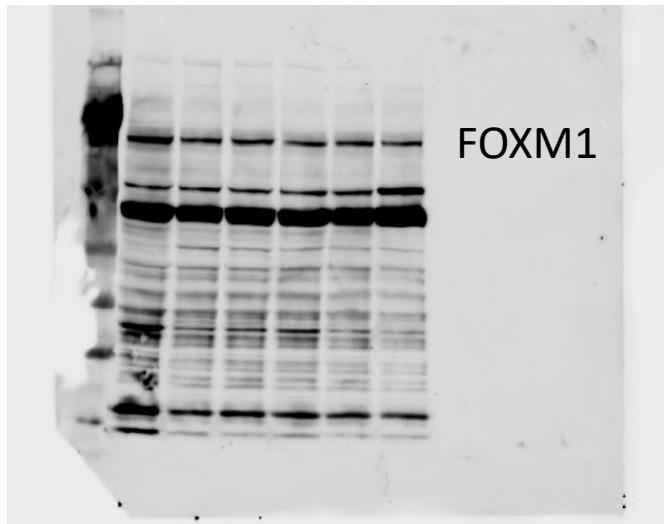


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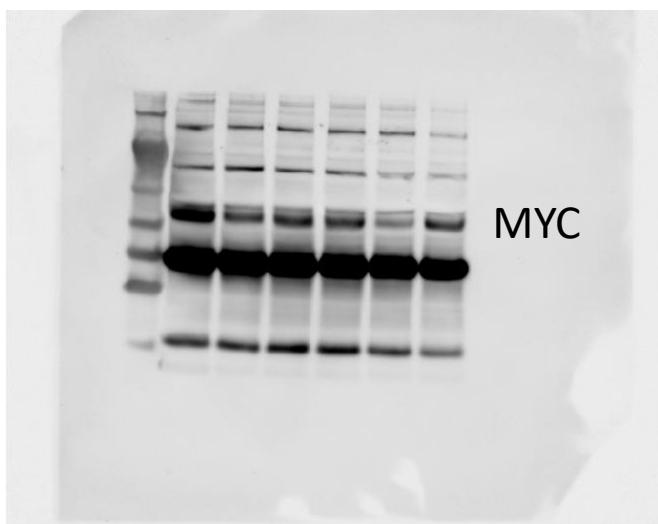


Lanes:
1) LNCaP
2) R1881
3) SARM-2F
4) T8039
5) GTx-024
6) Mitomycin-C

1 2 3 4 5 6

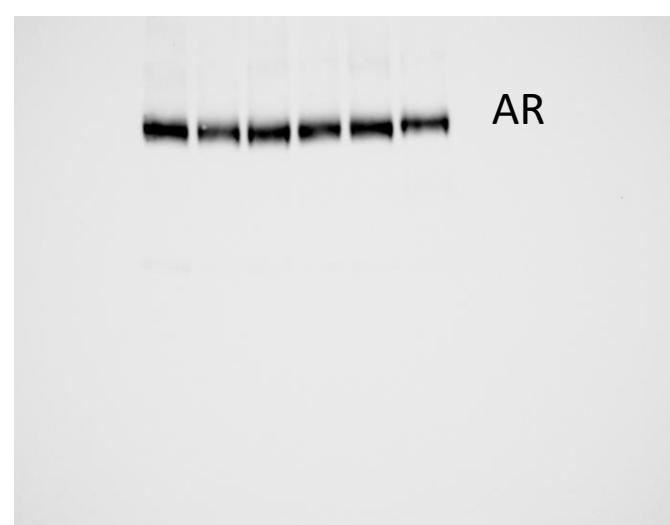


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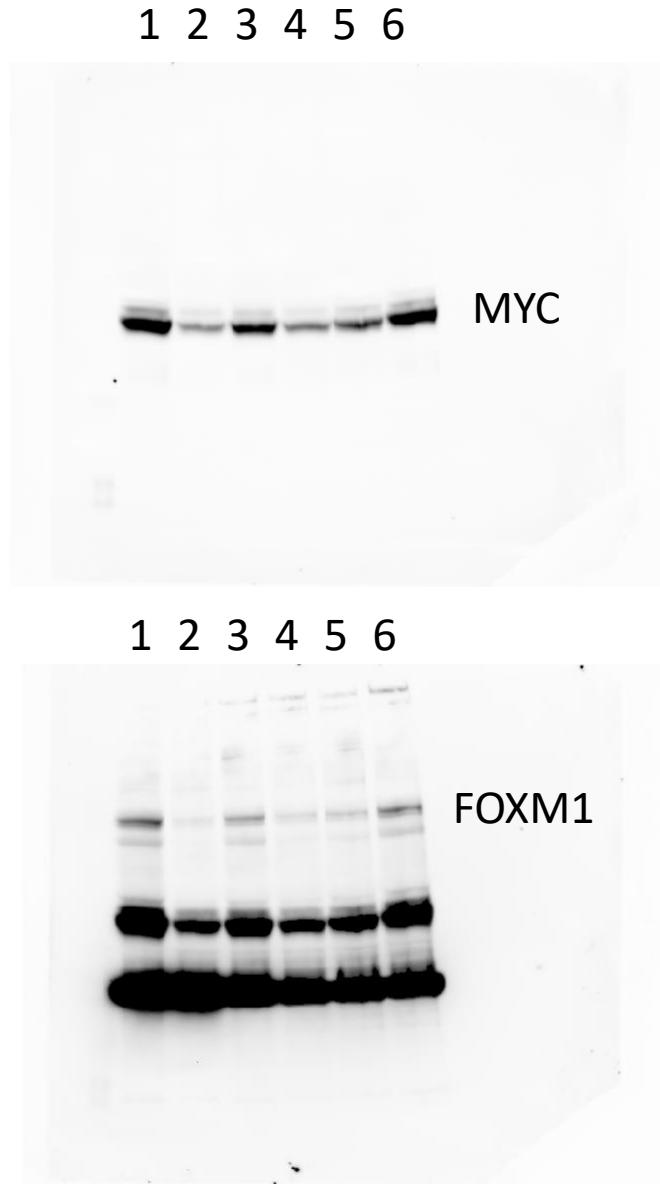
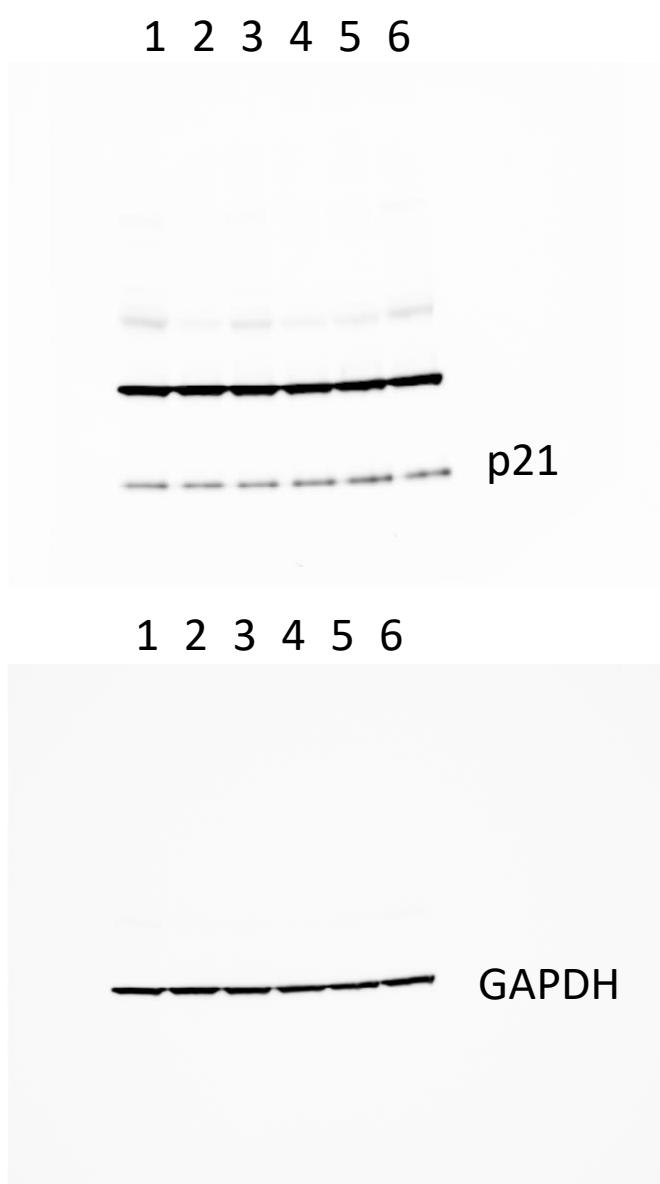


Lanes:
1) LNCaP
2) R1881
3) SARM-2F
4) T8039
5) GTx-024
6) Mitomycin-C

1 2 3 4 5 6



Full unedited gels for Figure 3B

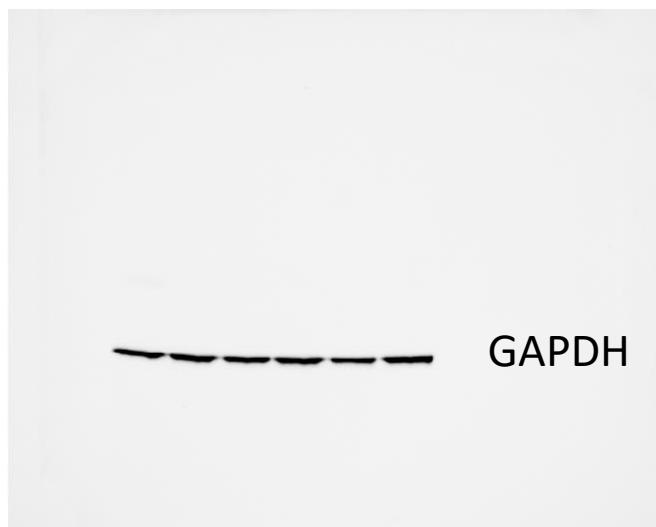


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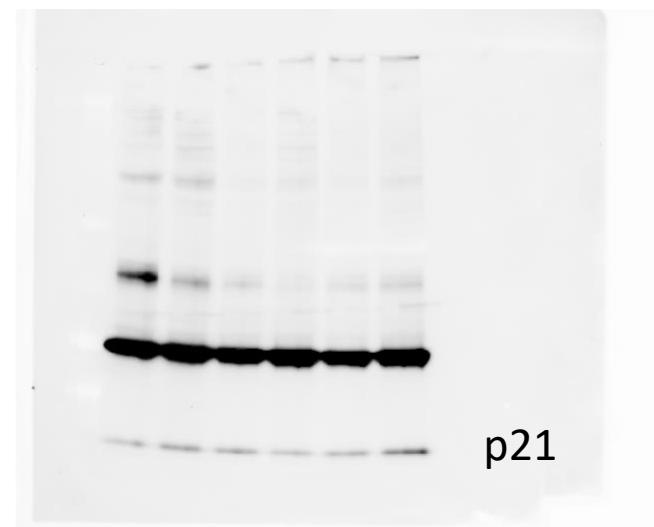
- 1) DMSO
- 2) R1881 1nM
- 3) ENZ 10uM
- 4) Cpd26 1uM
- 5) Cpd26 1nM
- 6) Cpd16 1pM

Full unedited gels for Figure 7G

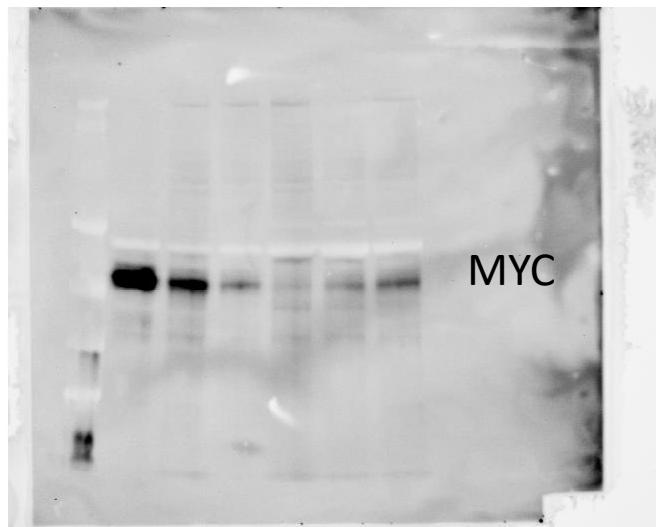
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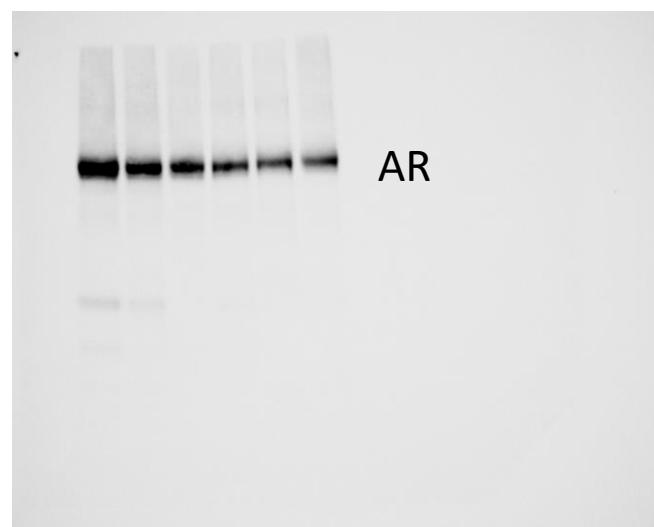
1 2 3 4 5 6



1 2 3 4 5 6



1 2 3 4 5 6



Lanes:

- 1) Vehicle
- 2) Testosterone
- 3) 15 mg/kg T8039
- 4) 30mg/kg T8039
- 5) 100mg/kg SARM-2F
- 6) 200mg/kg SARM-2F